<110>   Aros Applied Biotechnology ApS

<120>   Classification of Cancer

<130>   69167(302423)

<140>   US 10/584,653
<141>   2006-06-27

<150>   PCT/DK04/000914
<151>   2004-12-23

<150>   PA 2004 01843
<151>   2004-11-26

<150>   PA 2004 00586
<151>   2004-04-07

<150>   PA 2004 00096
<151>   2004-01-24

<150>   PA 2003 01940
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<160>   139

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<213>  NM_004184.3| Homo sapiens tryptophanyl-tRNA synthetase (WARS), transcript
variant 1, mRNA


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ggcccagatg ccacagaagc tgaagaggat tttgtggacc catggacagt acagacaagc    660

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transcript variant 1, mRNA


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<211> 3799

<212> DNA

<213> NM_003488.2 Homo sapiens A kinase (PRKA) anchor protein 1 (AKAP1), nuclear gene encoding mitochondrial protein, transcript variant 1, mRNA

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subunit 2 (PA28 beta) (PSME2), mRNA


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molecule 5 (CEACAM5), mRNA


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<211> 2033

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<210> 14

<211> 4180

<212> DNA

<213> NM_006291.2| Homo sapiens tumor necrosis factor, alpha-induced protein 2 (TNFAIP2), mRNA

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<210>  15

<211>  2524

<212>  DNA

<213>  NM_000249.2| Homo sapiens mutL homolog 1, colon cancer, nonpolyposis type
2 (E. coli) (MLH1), mRNA


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gcggggaag ttatccagcg gccagctaat gctatcaaag agatgattga gaactgttta      180

gatgcaaaat ccacaagtat tcaagtgatt gttaaagagg gaggcctgaa gttgattcag     240

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gatggaaagt gtgcatacag agcaagttac tcagatggaa aactgaaagc ccctcctaaa    480

ccatgtgctg gcaatcaagg gacccagatc acggtggagg accttttta caacatagcc     540

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ggcaggtatt cagtacacaa tgcaggcatt agtttctcag ttaaaaaaca aggagagaca    660

gtagctgatg ttaggacact acccaatgcc tcaaccgtgg acaatattcg ctccatcttt    720

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cagaatgtgg atgttaatgt gcaccccaca aagcatgaag ttcacttcct gcacgaggag   1020

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ataa   2524

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<211> 1536

<212> DNA

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gacaccatca aaaccaaccc tgacgacaga agaatcatca tgtgcgcttg gaatccaaga   660

gatcttcctc tgatggcgct gcctccatgc catgccctct gccagttcta tgtggtgaac   720

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aaaattcagc ttcagcgaga acccagacct ttcccaaagc tcaggattct tcgaaaagtt      960

gagaaaattg atgacttcaa agctgaagac tttcagattg aagggtacaa tccgcatcca     1020

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<210>  17

<211>  2986

<212>  DNA

<213>  NM_000201.1| Homo sapiens intercellular adhesion molecule 1 (CD54), human
rhinovirus receptor (ICAM1), mRNA


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<211> 736

<212> DNA

<213> NM_004492.1| Homo sapiens general transcription factor IIA, 2 (12kD subunit) (GTF2A2), mRNA

<400> 18
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<213>  NM_004850.3| Homo sapiens Rho-associated, coiled-coil containing protein
kinase 2 (ROCK2), mRNA


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<211>  1276

<212>  DNA

<213>  NM_003581.1| Homo sapiens NCK adaptor protein 2 (NCK2), mRNA


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<211>  1577

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<213>  NM_006214.2| Homo sapiens phytanoyl-CoA hydroxylase (Refsum disease)
(PHYH), mRNA


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<211> 3060

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<213> NM_004739.2| Homo sapiens metastais-associated gene family, member 2
(MTA2), mRNA

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<211> 2407

<212> DNA

<213> NM_001091.1| Homo sapiens amiloride binding protein 1 (amine oxidase
(copper-containing)) (ABP1), mRNA

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<211> 1094

<212> DNA

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variant 1, mRNA

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aagaatggag ttcaaacatg tctaaaccca gattcagcag atgtgaagga actgattaaa      300

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aagaaagttc tgaaagttcg aaaatctcaa cgttctcgtc aaaagaagac tacataagag      420

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ttccaaagga ggatggcata taatacaaag gcttattaat ttgactagaa aatttaaaac      540

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mRNA


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agcagcagga ggagaccggc gctgccaccc tgctactgca gggtgaggaa gaaggggaag      1020

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<210> 29

<211> 1575

<212> DNA

<213> NM_014298.3| Homo sapiens quinolinate phosphoribosyltransferase
(nicotinate-nucleotide pyrophosphorylase (carboxylating)) (QPRT), mRNA

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<211> 768

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<210> 31

<211> 696

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<213> NM_002984.1| Homo sapiens chemokine (C-C motif) ligand 4 (CCL4), mRNA

<400> 31
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ccataataaa actttaaaat aaaatgcaga cagtta                              696
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<210> 32

<211> 3338

<212> DNA

<213> NM_001455.2| Homo sapiens forkhead box O3A (FOXO3A), transcript variant 1, mRNA

<400> 32
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tgcgccaggt tcgctggccg cacgtcttca ggtcctcctg ttcctgggag gcgggcgcgg     120

caggactggg aggtggcggc agcggggcgg gactcgcccga ggacggggct ccggcccggg     180

ataaccaact ctccttctct cttctttggt gcttccccag gcggcggcgg cggcgcccgg     240

gagccggagc cttcgcggcg tccacgtccc tcccccgctg caccccgccc cggcgcgaga     300

ggagagcgcg agagccccag ccgcgggcgg gcgggcggcg aagatggcag aggcaccggc     360
```

```
ttccccggcc ccgctctctc cgctcgaagt ggagctggac ccggagttcg agccccagag      420

ccgtccgcga tcctgtacgt ggcccctgca aaggccggag ctccaagcga gccctgccaa      480

gccctcgggg gagacggccg ccgactccat gatccccgag gaggaggacg atgaagacga      540

cgaggacggc gggggacggg ccggctcggc catggcgatc ggcggcggcg gcgggagcgg      600

cacgctgggc tccgggctgc tccttgagga ctcggcccgg gtgctggcac ccggagggca      660

agaccccggg tctgggccag ccaccgcggc gggcgggctg agcgggggta cacaggcgct      720

gctgcagcct cagcaaccgc tgccaccgcc gcagccgggg gcggctgggg gctccgggca      780

gccgaggaaa tgttcgtcgc ggcggaacgc ctggggaaac ctgtcctacg cggacctgat      840

caccccgcgc atcgagagct ccccggacaa acggctcact ctgtcccaga tctacgagtg      900

gatggtgcgt tgcgtgccct acttcaagga taagggcgac agcaacagct ctgccggctg      960

gaagaactcc atccggcaca acctgtcact gcatagtcga ttcatgcggg tccagaatga     1020

gggaactggc aagagctctt ggtggatcat caaccctgat ggggggaaga gcggaaaagc     1080

ccccccggcg cgggctgtgt ctccatggac aa tagcaacaag tataccaaga gccgtggccg     1140

cgcagccaag aagaaggcag ccctgcagac agcccccgaa tcagctgacg acagtccctc     1200

ccagctctcc aagtggcctg gcagccccac gtcacgcagc agtgatgagc tggatgcgtg     1260

gacggacttc cgttcacgca ccaattctaa cgccagcaca gtcagtggcc gcctgtcgcc     1320

catcatggca agcacagagt ggatgaagt ccaggacgat gatgcgcctc tctcgcccat     1380

gctctacagc agctcagcca gcctgtcacc ttcagtaagc aagccgtgca cggtggaact     1440

gccacggctg actgatatgg caggcaccat gaatctgaat gatgggctga ctgaaaacct     1500

catggacgac ctgctggata acatcacgct cccgccatcc cagccatcgc ccactggggg     1560

actcatgcag cggagctcta gcttcccgta taccaccaag ggctcgggcc tgggctcccc     1620

aaccagctcc tttaacagca cggtgttcgg accttcatct ctgaactccc tacgccagtc     1680

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taaccagaca ctccaggacc tgctcacttc ggactcactt agccacagcg atgtcatgat     1800

gacacagtcg gaccccttga tgtctcaggc cagcaccgct gtgtctgccc agaattcccg     1860

ccggaacgtg atgcttcgca atgatccgat gatgtccttt gctgcccagc ctaaccaggg     1920

aagtttggtc aatcagaact tgctccacca ccagcaccaa acccagggcg ctcttggtgg     1980

cagccgtgcc ttgtcgaatt ctgtcagcaa catgggcttg agtgagtcca gcagccttgg     2040
```

```
gtcagccaaa caccagcagc agtctcctgt cagccagtct atgcaaaccc tctcggactc    2100

tctctcaggc tcctccttgt actcaactag tgcaaacctg cccgtcatgg gccatgagaa    2160

gttcccagc gacttggacc tggacatgtt caatgggagc ttggaatgtg acatggagtc    2220

cattatccgt agtgaactca tggatgctga tgggttggat tttaactttg attccctcat    2280

ctccacacag aatgttgttg gtttgaacgt ggggaacttc actggtgcta agcaggcctc    2340

atctcagagc tgggtgccag gctgaaggat cactgaggaa ggggaagtgg gcaaagcaga    2400

ccctcaaact gacacaagac ctacagagaa aacccttttgc caaatctgct ctcagcaagt    2460

ggacagtgat accgtttaca gcttaacacc tttgtgaatc ccacgccatt ttcctaaccc    2520

agcagagact gttaatggcc ccttaccctg ggtgaagcac ttaccttgg aacagaactc    2580

taaaaagtat gcaaaatctt ccttgtacag ggtggtgagc cgcctgccag tggaggacag    2640

caccctcag caccacccac cctcattcag agcacaccgt gagcccccgt cggccattct    2700

gtggtgtttt aatattgcga tggtttatgg gacgtttta gtgttgttct tgtgtttgtt    2760

ttcctttgac tttctgagtt tttcacatgc attaacttgc ggtatttttc tgttaaaatg    2820

ttaaccgtcc ttcccctagc aaatttaaaa acagaaagaa aatgttgtac cagttaccat    2880

tccgggttcg agcatcacaa gcttttgagc gcatggaact ccataaacta acaaattaca    2940

taaactaaag ggggattttc tttcttcttt tgtttggtag aaaattatcc ttttctaaaa    3000

actgaacaat ggcacaattg tttgctatgt gcacccgtcc aggacagaac cgtgcatagg    3060

caaaaggagt ggagcacagc gtccggccca gtgtgtttcc ggttctgagt cagggtgatc    3120

tgtggacggg accccagcac caagtctacg ggtgccagat cagtagggcc tgtgatttcc    3180

tgtcagtgtc ctcagctaat gtgaacagtg ttggtctgct ggttagaaac tagaatattg    3240

atattttcag gaaagaaatc agctcagctc tccactcatt gccaaatgtc actaaagggt    3300

ttagttttaa ggagaaagaa aaggaaaaaa aaaaaaaa                            3338
```

<210>   33

<211>   2646

<212>   DNA

<213>   NM_152873.1| Homo sapiens tumor necrosis factor receptor superfamily,
member 6 (TNFRSF6), transcript variant 4, mRNA

<400>   33

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ccagcgaggc ttccttccca tcctcctgac caccggggct tttcgtgagc tcgtctctga    120

tctcgcgcaa gagtgacaca caggtgttca aagacgcttc tggggagtga gggaagcggt    180

ttacgagtga cttggctgga gcctcagggg cgggcactgg cacggaacac accctgaggc    240

cagccctggc tgcccaggcg gagctgcctc ttctccccgcg ggttggtgga cccgctcagt    300

acggagttgg ggaagctctt tcacttcgga ggattgctca acaaccatgc tgggcatctg    360

gaccctccta cctctggttc ttacgtctgt tgctagatta tcgtccaaaa gtgttaatgc    420

ccaagtgact gacatcaact ccaagggatt ggaattgagg aagactgtta ctacagttga    480

gactcagaac ttggaaggcc tgcatcatga tggccaattc tgccataagc cctgtcctcc    540

aggtgaaagg aaagctaggg actgcacagt caatgggat gaaccagact gcgtgccctg    600

ccaagaaggg aaggagtaca cagacaaagc ccattttct tccaaatgca gaagatgtag    660

attgtgtgat gaaggacatg atgtgaacat ggaatcatca aggaatgcac actcaccagc    720

aacaccaagt gcaaagagga aggatccgaa tctaacttgg ggtggctttg tcttcttctt    780

ttgccaattc cactaattgt ttgggtgaag agaaaggaag tacagaaaac atgcagaaag    840

cacagaaagg aaaaccaagg ttctcatgaa tctccaacct taaatcctga aacagtggca    900

ataaatttat ctgatgttga cttgagtaaa tatatcacca ctattgctgg agtcatgaca    960

ctaagtcaag ttaaaggctt tgttcgaaag aatggtgtca atgaagccaa aatagatgag   1020

atcaagaatg acaatgtcca agacacagca gaacagaaag ttcaactgct tcgtaattgg   1080

catcaacttc atggaaagaa agaagcgtat gacacattga ttaaagatct caaaaaagcc   1140

aatctttgta ctcttgcaga gaaaattcag actatcatcc tcaaggacat tactagtgac   1200

tcagaaaatt caaacttcag aaatgaaatc caaagcttgg tctagagtga aaaacaacaa   1260

attcagttct gagtatatgc aattagtgtt tgaaaagatt cttaatagct ggctgtaaat   1320

actgcttggt tttttactgg gtacatttta tcatttatta gcgctgaaga gccaacatat   1380

ttgtagattt ttaatatctc atgattctgc ctccaaggat gtttaaaatc tagttgggaa   1440

aacaaacttc atcaagagta aatgcagtgg catgcctaagt acccaaatag gagtgtatgc   1500

agaggatgaa agattaagat tatgctctgg catctcaact atgattctgt agtatgaatg   1560

taatcagtgt atgttagtac aaatgtctat ccacaggcta accccactct atgaatcaat   1620

agaagaagct atgacctttt gctgaaatat cagttactga acaggcaggc cactttgcct   1680

ctaaattacc tctgataatt ctagagattt taccatattt ctaaactttg tttataactc   1740
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tgagaagatc atatttatgt aaagtatatg tatttgagtg cagaatttaa ataaggctct    1800

acctcaaaga cctttgcaca gtttattggt gtcatattat acaatatttc aattgtgaat    1860

tcacatagaa aacattaaat tataatgttt gactattata tatgtgtatg cattttactg    1920

gctcaaaact acctacttct ttctcaggca tcaaaagcat tttgagcagg agagtattac    1980

tagagctttg ccacctctcc atttttgcct tggtgctcat cttaatggcc taatgcaccc    2040

ccaaacatgg aaatatcacc aaaaaaatact taatagtcca ccaaaaggca agactgccct    2100

tagaaattct agcctggttt ggagatacta actgctctca gagaaagtag ctttgtgaca    2160

tgtcatgaac ccatgtttgc aatcaaagat gataaaatag attcttattt ttcccccacc    2220

cccgaaaatg ttcaataatg tcccatgtaa aacctgctac aaatggcagc ttatacatag    2280

caatggtaaa atcatcatct ggatttagga attgctcttg tcataccccc aagtttctaa    2340

gatttaagat tctccttact actatcctac gtttaaatat ctttgaaagt ttgtattaaa    2400

tgtgaatttt aagaaataat atttatattt ctgtaaatgt aaactgtgaa gatagttata    2460

aactgaagca gatacctgga accacctaaa gaacttccat ttatggagga ttttttttgcc    2520

ccttgtgttt ggaattataa aatataggta aaagtacgta attaaataat gttttttggta    2580

aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa    2640

aaaaaa                                                               2646
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<210> 34

<211> 817

<212> DNA

<213> NM_002038.2| Homo sapiens interferon, alpha-inducible protein (clone IFI-6-16) (G1P3), transcript variant 1, mRNA

<400> 34

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cttctctcct ccaaggtcta gtgacggagc ccgcgcgcgg cgccaccatg cggcagaagg     120

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agaaaaagtg ctcggagagc tcggacagcg gctccgggtt ctggaaggcc ctgaccttca     240

tggccgtcgg aggaggactc gcagtcgccg ggctgcccgc gctgggcttc accggcgccg     300

gcatcgcggc caactcggtg gctgcctcgc tgatgagctg gtctgcgatc ctgaatgggg     360
```

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gcggcgtgcc cgccggggggg ctagtggcca cgctgcagag cctcgggggct ggtggcagca      420

gcgtcgtcat aggtaatatt ggtgccctga tgggctacgc cacccacaag tatctcgata      480

gtgaggagga tgaggagtag ccagcagctc ccagaacctc ttcttccttc ttggcctaac      540

tcttccagtt aggatctaga actttgcctt tttttttttt tttttttttt tttgagatgg      600

gttctcacta tattgtccag gctagagtgc agtggctatt cacagatgcg aacatagtac      660

actgcagcct ccaactccta gcctcaagtg atcctcctgt ctcaacctcc caagtaggat      720

tacaagcatg cgccgacgat gcccagaatc cagaactttg tctatcactc tccccaacaa      780

cctagatgtg aaaacagaat aaacttcacc cagaaaa                                817
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<400>   35
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attcaaggag tacctctctc tagaaccgta cgctgtacct gcatcagcat tagtaatcaa      180

cctgttaatc caaggtcttt agaaaaactt gaaattattc ctgcaagcca attttgtcca      240

cgtgttgaga tcattgctac aatgaaaaag aagggtgaga agagatgtct gaatccagaa      300

tcgaaggcca tcaagaatt actgaaagca gttagcaagg aaatgtctaa aagatctcct      360

taaaaccaga ggggagcaaa atcgatgcag tgcttccaag gatggaccac acagaggctg      420

cctctcccat cacttcccta catggagtat atgtcaagcc ataattgttc ttagtttgca      480

gttacactaa aaggtgacca atgatggtca ccaaatcagc tgctactact cctgtaggaa      540

ggttaatgtt catcatccta gctattcag taataactct accctggcac tataatgtaa      600

gctctactga ggtgctatgt tcttagtgga tgttctgacc ctgcttcaaa tatttcctc      660

accttttccca tcttccaagg gtactaagga atctttctgc tttgggggttt atcagaattc      720

tcagaatctc aaataactaa aaggtatgca atcaaatctg cttttttaaag aatgctctttt      780

acttcatgga cttccactgc catcctccca aggggccaa attctttcag tggctaccta      840
```

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catacaattc caaacacata caggaaggta gaaatatctg aaaatgtatg tgtaagtatt      900

cttatttaat gaaagactgt acaaagtata agtcttagat gtatatattt cctatattgt      960

tttcagtgta catggaataa catgtaatta agtactatgt atcaatgagt aacaggaaaa     1020

ttttaaaaat acagatagat atatgctctg catgttacat aagatataatg tgctgaatgg     1080

ttttcaaata aaaatgaggt actctcctgg aaatattaag aaagactatc taaatgttga     1140

aagatcaaaa ggttaataaa gtaattataa ct                                  1172


<210>  36

<211>  396

<212>  DNA

<213>  NM_005950.1┤ Homo sapiens metallothionein 1G (MT1G), mRNA


<400>  36
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cctcgggttg caatggaccc caactgctcc tgtgccgctg gtgtctcctg cacctgcgcc      120

agctcctgca agtgcaaaga gtgcaaatgc acctcctgca agaagagctg ctgctcctgc      180

tgccctgtgg gctgtgtgccaa gtgtgcccaa ggctgcatct gcaaagggggc atcggagaag      240

tgcagctgct gcgcctgatg tcgggacagc cctgctccca agtacaaata gagtgacccg      300

taaaatctag gattttttgt tttttgctac aatcttgacc cctttgctac attcccttt      360

ttctgtgaaa tatgtgaata ataattaaac acttag                              396


<210>  37

<211>  2755

<212>  DNA

<213>  NM_000043.3┤ Homo sapiens tumor necrosis factor receptor superfamily,
member 6 (TNFRSF6), transcript variant 1, mRNA


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ccagcgaggc ttccttccca tcctcctgac caccggggct tttcgtgagc tcgtctctga      120

tctcgcgcaa gagtgacaca caggtgttca aagacgcttc tggggagtga gggaagcggt      180
```

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ttacgagtga cttggctgga gcctcagggg cgggcactgg cacggaacac accctgaggc    240

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gaccctccta cctctggttc ttacgtctgt tgctagatta tcgtccaaaa gtgttaatgc    420

ccaagtgact gacatcaact ccaagggatt ggaattgagg aagactgtta ctacagttga    480

gactcagaac ttggaaggcc tgcatcatga tggccaattc tgccataagc cctgtcctcc    540

aggtgaaagg aaagctaggg actgcacagt caatggggat gaaccagact gcgtgccctg    600

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attgtgtgat gaaggacatg gcttagaagt ggaaataaac tgcacccgga cccagaatac    720

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taaaggcttt gttcgaaaga atggtgtcaa tgaagccaaa atagatgaga tcaagaatga    1140

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tggaaagaaa gaagcgtatg acacattgat taaagatctc aaaaaagcca atctttgtac    1260

tcttgcagag aaaattcaga ctatcatcct caaggacatt actagtgact cagaaaattc    1320

aaacttcaga aatgaaatcc aaagcttggt ctagagtgaa aaacaacaaa ttcagttctg    1380

agtatatgca attagtgttt gaaaagattc ttaatagctg gctgtaaata ctgcttggtt    1440

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gattaagatt atgctctggc atctaacata tgattctgta gtatgaatgt aatcagtgta    1680

tgttagtaca aatgtctatc cacaggctaa ccccactcta tgaatcaata gaagaagcta    1740

tgaccttttg ctgaaatatc agttactgaa caggcaggcc actttgcctc taaattacct    1800

ctgataattc tagagatttt accatatttc taaactttgt ttataactct gagaagatca    1860
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tatttatgta aagtatatgt atttgagtgc agaatttaaa taaggctcta cctcaaagac    1920

ctttgcacag tttattggtg tcatattata caatatttca attgtgaatt cacatagaaa    1980

acattaaatt ataatgtttg actattatat atgtgtatgc attttactgg ctcaaaacta    2040

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cacctctcca tttttgcctt ggtgctcatc ttaatggcct aatgcacccc caaacatgga    2160

aatatcacca aaaaatactt aatagtccac caaaaggcaa gactgccctt agaaattcta    2220

gcctggtttg gagatactaa ctgctctcag agaaagtagc tttgtgacat gtcatgaacc    2280

catgtttgca atcaaagatg ataaaataga ttcttatttt tcccccaccc ccgaaaatgt    2340

tcaataatgt cccatgtaaa acctgctaca aatggcagct tatacatagc aatggtaaaa    2400

tcatcatctg gatttaggaa ttgctcttgt cataccccca agtttctaag atttaagatt    2460

ctccttacta ctatcctacg tttaaatatc tttgaaagtt tgtattaaat gtgaatttta    2520

agaaataata tttatatttc tgtaaatgta aactgtgaag atagttataa actgaagcag    2580

atacctggaa ccacctaaag aacttccatt tatggaggat tttttgccc cttgtgtttg      2640

gaattataaa atataggtaa aagtacgtaa ttaaataatg tttttggtaa aaaaaaaaaa    2700

aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaa          2755


<210>  38

<211>  1600

<212>  DNA

<213>  NM_001953.2| Homo sapiens endothelial cell growth factor 1 (platelet-
derived) (ECGF1), mRNA


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gggcgaggcc gggtacctgg gctgggatcc ggagcaagcg ggcgagggca gcgccctaag     120

caggcccgga gcgatggcac ccttgatgac cccgggaacc ggggcgcccac ccgcgcctgg    180

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cttcgtggcc gctgtggtga atgggagcgc gcagggcgca cagatcgggg ccatgctgat    360

ggccatccga cttcgggggca tggatctgga ggagacctcg gtgctgaccc aggccctggc    420
```

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tcagtcggga cagcagctgg agtggccaga ggcctggcgc cagcagcttg tggacaagca    480

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taagctggag tctattcctg gattcaatgt catccagagc ccagagcaga tgcaagtgct    660

gctggaccag gcgggctgct gtatcgtggg tcagagtgag cagctggttc ctgcggacgg    720

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ggaggcgctc gtactctccg accgcgcgcc attcgccgcc ccctcgccct tcgcagagct   1560

cgttctgccg ccgcagcaat aaagctcctt tgccgcgaaa                          1600
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<210> 39

<211> 931

<212> DNA

<213> NM_005138.1| Homo sapiens SCO cytochrome oxidase deficient homolog 2 (yeast) (SCO2), nuclear gene encoding mitochondrial protein, mRNA

<400> 39
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caggctctct cagctcaagc ctccggtcct ccctgggacc ctgggaggcc aggccctgca      180

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tggggcctgg ctggccctga gggctgagaa ggagaggctg cagcagcaaa agcgaacaga      360

agccctgcgc caggcagctg tgggccaggg cgacttccac ctgctggatc acagaggccg      420

ggctcgctgc aaggctgact tccggggcca gtgggtgctg atgtactttg gcttcactca      480

ctgccctgac atctgcccag acgagctgga gaagctggtg caggtggtgc ggcagctgga      540

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cgacgttgaa gccatggccc gctacgtcca ggacttccac ccaagactgt tgggtctgac      660

cggctccacc aaacaggttg cccaggctag tcacagttac cgcgtgtact acaatgccgg      720

ccccaaggat gaggaccagg actacatcgt ggaccactcc attgccatct acctgctcaa      780

ccctgacggc ctcttcacgg attactacgg ccggagcaga tcggctgagc agatctcaga      840

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ggccccatca ttaaacgggc tgcgtttaaa a                                     931


<210>  40

<211>  1216

<212>  DNA

<213>  NM_006419.1| Homo sapiens chemokine (C-X-C motif) ligand 13 (B-cell
chemoattractant) (CXCL13), mRNA


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gctcaagtct gaactctacc tccagacaga atgaagttca tctcgacatc tctgcttctc      120

atgctgctgg tcagcagcct ctctccagtc caaggtgttc tggaggtcta ttacacaagc      180

ttgaggtgta gatgtgtcca agagagctca gtctttatcc ctagacgctt cattgatcga      240

attcaaatct tgcccgtgg gaatggttgt ccaagaaaag aaatcatagt ctggaagaag      300

aacaagtcaa ttgtgtgtgt ggaccctcaa gctgaatgga tacaaagaat gatggaagta      360

ttgagaaaaa gaagttcttc aactctacca gttccagtgt ttaagagaaa gattccctga      420

tgctgatatt tccactaaga acacctgcat tcttcccta tccctgctct ggattttagt      480
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ggactatgtg aaaaataacc ttgcggagc tgatgggca aactcaagct tcttcactca      600

cagcacccta tatacacttg gagtttgcat tcttattcat cagggaggaa agtttctttg      660

aaaatagtta ttcagttata agtaatacag gattattttg attatatact tgttgtttaa      720

tgtttaaaat ttcttagaaa acaatggaat gagaatttaa gcctcaaatt tgaacatgtg      780

gcttgaatta agaagaaaat tatggcatat attaaaagca ggcttctatg aaagactcaa      840

aaagctgcct gggaggcaga tggaacttga gcctgtcaag aggcaaagga atccatgtag      900

tagatatcct ctgcttaaaa actcactacg gaggagaatt aagtcctact tttaaagaat      960

ttctttataa aatttactgt ctaagattaa tagcattcga agatccccag acttcataga      1020

atactcaggg aaagcattta aagggtgatg tacacatgta tcctttcaca catttgcctt      1080

gacaaacttc tttcactcac atctttttca ctgacttttt ttgtgggggc ggggccgggg      1140

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agcaaacatt ttactt                                                      1216


<210>  41

<211>  738

<212>  DNA

<213>  NM_006433.2  Homo sapiens granulysin (GNLY), transcript variant NKG5,
mRNA


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aaagattaag ctgcaggctc cctgcccata aaacagggtg tgaaaggcat ctcagcggct      120

gccccaccat ggctacctgg gccctcctgc tccttgcagc catgctcctg ggcaacccag      180

gtctggtctt ctctcgtctg agccctgagt actacgacct ggcaagagcc cacctgcgtg      240

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agatggtgga taagcccacc cagagaagtg tttccaatgc tgcgaccogg gtgtgtagga      420

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ttacccaggg cctcgtggcc ggagaaactg cccagcagat ctgtgaggac ctcaggttgt      540
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ctcctgtcct cagatcccgg gaacctcagc aacctctgcc ggctcctcgc ttcctcgatc      660

cagaatccac tctccagtct ccctcccctg actccctctg ctgtcctccc ctctcacgag      720

aataaagtgt caagcaag                                                    738


<210>   42

<211>   1579

<212>   DNA

<213>   NM_001767.2| Homo sapiens CD2 antigen (p50), sheep red blood cell
receptor (CD2), mRNA


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tgtttcttcc aaaggtgcag tctccaaaga gattacgaat gccttggaaa cctggggtgc      120

cttgggtcag gacatcaact tggacattcc tagttttcaa atgagtgatg atattgacga      180

tataaaatgg gaaaaaactt cagacaagaa aaagattgca caattcagaa aagagaaaga      240

gactttcaag gaaaaagata catataagct atttaaaaat ggaactctga aaattaagca      300

tctgaagacc gatgatcagg atatctacaa ggtatcaata tatgatacaa aaggaaaaaa      360

tgtgttggaa aaaatatttg atttgaagat tcaagagagg gtctcaaaac caaagatctc      420

ctggacttgt atcaacacaa ccctgacctg tgaggtaatg aatggaactg accccgaatt      480

aaacctgtat caagatggga aacatctaaa actttctcag agggtcatca cacacaagtg      540

gaccaccagc ctgagtgcaa aattcaagtg cacagcaggg aacaaagtca gcaaggaatc      600

cagtgtcgag cctgtcagct gtccagagaa aggtctggac atctatctca tcattggcat      660

atgtggagga ggcagcctct tgatggtctt tgtggcactg ctcgttttct atattaccaa      720

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agcaacttcc caacatcctc ctccaccacc tggtcatcgt tcccaggcac ctagtcatcg      900

tcccccgcct cctggacacc gtgttcagca ccagcctcag aagaggcctc ctgctccgtc      960

gggcacacaa gttcaccagc agaaaggccc gccctcccc agacctcgag ttcagccaaa      1020

acctccccat ggggcagcag aaaactcatt gtcccccttcc tctaattaaa aagagatagaa      1080
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actgtctttt tcaataaaaa gcactgtgga tttctgccct cctgatgtgc atatccgtac      1140

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accgagcaca gaaatcttag agatttcttg tcccctctca ggtcatgtgt agatgcgata      1380

aatcaagtga ttggtgtgcc tgggtctcac tacaagcagc ctatctgctt aagagactct      1440

ggagtttctt atgtgccctg gtggacactt gcccaccatc ctgtgagtaa aagtgaaata      1500

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aaaaaaaaaa aaaaaaaaa                                                    1579


<210>   43

<211>   3738

<212>   DNA

<213>   NM_006275.4| Homo sapiens splicing factor, arginine/serine-rich 6
(SFRS6), mRNA


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tgttcgggct cttgccgtcc ccgcacccgc accgcggtta ctggcttgcg gtccgccgtt      120

cgacaaccag cccttgggtc cccgcccgcc acggacatgc cgcgcgtcta cataggacgc      180

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cacgcccggg gcccgcgtcg cgatcgcgac ggctacagct acggaagccg cagtggtgga      420

ggtggataca gcagtcggag aacatctgac agagacaaat acggaccacc tgttcgtaca      480

gaatacaggc ttattgtaga aaatctttct agtcggtgca gttggcaaga tttaaaggat      540

tttatgcgac aagcaggtga agtaacctat gcggatgccc acaaggaacg aacaaatgag      600

ggtgtaattg agtttcgctc ctactctgac atgaagcgtg ctttggacaa actggatggc      660

acagaaataa atggcagaaa tattaggctt attgaagata agccacgcac aagccatagg      720

cgatcttact ctggaagcag atccaggtct cgatctagaa gacggtcacg aagtaggagt      780
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tatgagaaat ctcgaagcag gtctcggtcc cgatcccca aagaaaatgg aaagggtgat    1020

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ccttctttgt agaattaaga taactttgat tttatagctt ttgagctaac gtaactttg    1440

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ctgttttgag ctttgaatac ttaaggcttt agaggggagaa cccaatttc aattatgttg   1620

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gtttggataa cattattttg tgcagttaat caaatttgcc aaagtcttta tctgcccctt   1860

taacaagttg agtaaaaata aaaggtattt tttagtcaat gtgttccatg attttgctta   1920

aattaatact tttaagtaat ggactttttt tcaaaggcaa atttaaacta tttaagaaat   1980

agctcctaat acttgggatc ttgtttagag aatccacttt ctggaagttc tcagcataat   2040

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aagatcatta tgtttaacag gcacttaatc tcagtaaagt cagttgccag ttaagttcca   2220

cccagtagtc agtcccttt gtagttagtg ggattatttg ataattggtt agatcatatt   2280

tgtaaatttt aatgctttgt gtaattggtt tgaaaaacag tgaaatgggt aaacgcaaaa   2340

cttttgtact ttattacgag taaagtgtaa tgagtactgt ggaaaccaaa tttgaatact   2400

gcaaatttgt aggagttact aggttagcaa ttagtccata catccataag cctgatgagt   2460

tgaaattgca gtttgagaag tgaattaacc ttacatccct ttgttcagat accttaaaag   2520
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agggttggtt tgtagattca aataatccag aaatatacct aataagattg agtgaaaaat    2760

ttgagtcaaa tatctagggc attcacagag tagctgtgag ttcttggtaa tgtgaaaaag    2820

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tagactggtt tgactatatt gttagctgcc acagtaagca ggtcattgta taggtaaatg    3300

cctgcaccca taattttcta gtaatagcca cgaccaattt attaacagtc agggcctatc    3360

cttgcctgta gttctcagtc actggatgca caaaatcact gtgtaacatt ggctcacttg    3420

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ctttatggcc agtttgagtc ctgcctactt tgacttttac gttcccattc ctgtgttacc    3660

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<211>   2033

<212>   DNA

<213>   NM_003212.1| Homo sapiens teratocarcinoma-derived growth factor 1
(TDGF1), mRNA


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ctgtttggc aatgactctg aattaaagcg atgctaacgc ctcttttccc cctaattgtt     240

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tccatctcgg ggatacctgg ccttcagaga tgacagcatt tggccccagg aggagcctgc     420

aattcggcct cggtcttccc agcgtgtgcc gcccatgggg atacagcaca gtaaggagct     480

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ccgctgcttt cctcaggcat ttctacccgg ctgtgatggc cttgtgatgg atgagcacct     720

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aacatttcct tacagaacta cttcttactt ctttgccctg ccctctccca aaaaactact     1020

tcttttttca aaagaaagtc agccatatct ccattgtgcc taagtccagt gtttcttttt     1080

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agtaactggg attacaggca tgtgtcacca tgcccagcta atttttttgt attttagtag     1260

agatggggggt ttcaccatat tggccagtct ggtctcgaac tctgaccttg tgatccatcg     1320

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ccagagtgct gaaggaatgg aagtccatct gcgtgtgtgc agggagactg ggtaggaaag     1620

aggaagcaaa tagaagagag aggttgaaaa acaaaatggg ttacttgatt ggtgattagg     1680

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aagctatata agacaagaac tccccttttt ttcccaaagg cattataaaa agaatgaagc    1800

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tcctccaagc tattcaattc tttttaactgt tgtagaagac aaaatgttca caatatattt    1920

agttgtaaac caagtgatca aactacatat tgtaaagccc attttttaaaa tacattgtat    1980

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<210>  45

<211>  367

<212>  DNA

<213>  NM_005951.1| Homo sapiens metallothionein 1H (MT1H), mRNA


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gcgcctgcgc cggctcctgc aagtgcaaaa agtgcaaatg cacctcctgc aagaagagct    120

gctgctcctg ttgccccctg ggctgtgcca agtgtgccca gggctgcatc tgcaaagggg    180

cgtcagagaa gtgcagctgc tgtgcctgat gtcggacag ccctgctgtc agatgaaaac     240

agaatgacac gtaaaatccg aggttttttt tttctacaac tccgactcat ttgctacatt    300

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aaaaaaa                                                              367


<210>  46

<211>  3052

<212>  DNA

<213>  NM_000767.4| Homo sapiens cytochrome P450, family 2, subfamily B,
polypeptide 6 (CYP2B6), mRNA


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ccttttggga aaccttctgc agatggatag aagaggccta ctcaaatcct ttctgaggtt    180

ccgagagaaa tatgggggacg tcttcacggt acacctggga ccgaggcccg tggtcatgct    240
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aaaccgctgg aaggtgcttc ggcgattctc tgtgaccact atgagggact tcgggatggg    420

aaagcggagt gtggaggagc ggattcagga ggaggctcag tgtctgatag aggagcttcg    480

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cagcgccccc aaggacctca tcgacacacta cctgctccaa atggaaaaag agaaatccaa    840

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ggcactgaaa aagactgaag cttttatccc cttctcctta gggaagcgga tttgtcttgg   1320

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caaaataccc ccaacatacc agatccgctt cctgccccgc tgaaggggct gagggaaggg   1500

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<210> 56

<211> 2090

<212> DNA

<213> NM_003242.3| Homo sapiens transforming growth factor, beta receptor II
(70/80kDa) (TGFBR2), mRNA

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<211> 4568

<212> DNA

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<400> 57

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<211> 2069

<212> DNA

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aatctttctc gcctactgcc tctctcgtgc cataacaaat aaccagtatg agatagtgta      840

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<210>  59

<211>  2402

<212>  DNA

<213>  NM_021200.1| Homo sapiens pleckstrin homology domain containing, family B
(evectins) member 1 (PLEKHB1), mRNA


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aa                                                                     2402


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<211>  2856

<212>  DNA

<213>  NM_003661.2| Homo sapiens apolipoprotein L, 1 (APOL1), transcript variant
1, mRNA


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ccttagctgg caatacttac caactcacac gaggcattgg gaaggacatc cgtgtgcctca      1020

gacgagccag agccaatctt cagtcagtac cgcatgcctc agcctcacgc ccccgggtca      1080

ctgagccaat ctcagctgaa agcggtgaac aggtggagag ggttaatgaa cccagcatcc      1140

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<210> 61

<211> 1655

<212> DNA

<213> NM_002164.3| Homo sapiens indoleamine-pyrrole 2,3 dioxygenase (INDO), mRNA

<400> 61

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aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaa                               1655


<210>  62

<211>  2242

<212>  DNA

<213>  NM_021784.3| Homo sapiens forkhead box A2 (FOXA2), transcript variant 1,
mRNA


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gcggctcggg caacatgagc gcgggctcca tgaacatgtc gtcgtacgtg ggcgctggca     420

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cggtgtaaaa ccatgtagtt ttaacagaac cagagggttg tactattgtt taaaaacagg   2040

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caatcttgac acggtgaaat ccaggtctcg ggtccgatta atttatggtt tctgcgtgct   2160

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<210> 63

<211> 1047

<212> DNA

<213> NM_033423.2| Homo sapiens granzyme H (cathepsin G-like 2, protein h-CCPX) (GZMH), mRNA

<400> 63

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gaggccaagc cccactcccg ccoctacatg gcctttgttc agtttctgca agagaagagt     180

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aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa      1020

aaaaaaaaaa aaaaaaaaaa aaaaaaa                                         1047


<210>   64

<211>   5243

<212>   DNA

<213>   NM_001165.3| Homo sapiens baculoviral IAP repeat-containing 3 (BIRC3),
transcript variant 1, mRNA


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aggaaaacga cttcttctag attttttttt cagtttcttc tataaatcaa aacatctcaa      180

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<211>  980
<212>  DNA
<213>  NM_003283.3| Homo sapiens troponin T1, skeletal, slow (TNNT1), mRNA


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tcggacaccg aggagcagga atatgaggag gagcagccgg aagaggaggc tgcggacgag       120

gaggaggaag cccccgaaga gccggagccg gtggcagagc cagaagagga acgccccaaa       180

ccaagccgcc ccgtggtgcc tcctttgatc ccgccaaaga tcccagaagg ggagcgcgtt       240

gacttcgatg acatccaccg caagcgcatg gagaaagacc tgctggagct gcagacactc       300

atcgatgtac atttcgagca gcggaagaag gaggaagagg agctggttgc cttgaaggag       360

cgcattgagc ggcgcggtc agagagagcc gagcaacagc gcttcagaac tgagaaggaa       420

cgcgaacgtc aggctaagct ggcggaggag aagatgagga aggaagaga agaggccaag       480

aagcgggcag aggatgatgc caagaaaaag aaggtgctgt ccaacatggg ggcccatttt      540

ggcggctacc tggtcaaggc agaacagaag cgtggtaagc ggcagacggg gcgggagatg      600
```

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aaggtgcgca tcctctccga gcgtaagaag cctctggaca ttgactacat gggggaggaa      660

cagctccggg cccggtctgc ctggctgcct ccatcacagc cctcctgccc tgccagggag      720

aaagcccagg agctgtcgga ctggatccac cagctggagt ctgagaagtt cgacctgatg      780

gcgaagctga aacagcagaa atatgagatc aacgtgctgt acaaccgcat cagccacgcc      840

cagaagttcc ggaaggggc aggga agggc cgcgttggag gccgctggaa gtgaggatgc      900

cgccccggac agtggcacct gggaagcctg ggagtgtttg tcccatcggt agcttgaaat      960

aaacgctccc ctcagacacc                                                  980


<210>  73
<211>  2213
<212>  DNA
<213>  NM_004067.1| Homo sapiens chimerin (chimaerin) 2 (CHN2), mRNA


<400>  73
gggcgtgcaa aggcgcggag cgggacggaa accacaaata aatagcggcg cgcggcagcgc       60

gtcatctggt ggagcaggaa gtgcaggcag agtccggagg ctggtgcttt ctgcgcgtcc      120

ccaggacttt gccatgggct gggggccgcg gaggctgcga gcggccgggc gagggcagcg      180

gcggcggcgt ccccaccggg gctgagcgag cagcgacgcg agggcgcgc ggagatggca      240

gcgtccagca actccagcct gtccggctcg tcggtgtcct ccgatgctga agaataccag      300

cctcctatat ggaaatcata cttatatcag ttacagcaag aggcacctcg tcccaagaga      360

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atcatctctc gggagcaggc ggatgagctt cttggaggcg tggagggtgc ctacatcctt      480

agagaaagcc agcggcaacc aggatgctac acgctggctc tcaggtttgg aaaccagacc      540

ttaaactaca ggctcttcca cgacgggaaa cactttgtgg gtgagaagag gtttgagtcg      600

attcatgatc tggtgacaga tggcttgata acactgtaca tagaaacaaa agctgccgag      660

tacatttcaa aaatgacaac taaccccatc tatgaacaca ttggatatgc caccctactc      720

agagaaaaag tatccagaag gctgagcagg tctaaaaatg aaccaagaaa aacaaacgtc      780

acacatgaag aacacacagc ggtggaaaag atctcctccc tggttcgaag ggctgccctc      840

acacacaacg acaaccactt caattatgag aagacacaca actttaaggt ccacacgttc      900

cgaggcccac actggtgtga atattgtgcc aatttcatgt gggggctcat cgcccaaggg      960
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gtccggtgct cagactgtgg attgaacgta cacaaacagt gttccaagca cgttcccaat     1020

gactgccaac ctgatctcaa gaggatcaag aaagtgtact gttgtgacct cacaacactt     1080

gtgaaggctc acaacactca gagacccatg gtggtagaca tatgcattcg ggaaattgaa     1140

gcaagaggat taaaatcgga aggcctttac agagtctctg ggttcactga cacacattgaa    1200

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gtcatcacat atgatacct atteccaattt atagatgcag caaaaatctc caatgcagat     1380

gagaggctgg aagccgtcca tgaagtgctg atgctgctgc ctcctgccca ctatgaaacc     1440

ctccggtacc taatgatcca cctcaaaaag gttactatga atgaaaaga caatttcatg     1500

aatgcagaaa atctgggat cgtgtttggg cccatctga tgaggccccc tgaggacagc     1560

accctgacca ccctgcatga tatgcggtac caaaagctga ttgtgcagat tttaatagaa     1620

aacgaagacg ttttattcta atccatcagg gaaatgagct gaatggccca gcaccatcaa     1680

gttgacacag ctaaggataa aacatttctt accacttgat ttgtttccca agcaagtgct    1740

agaatttgct ggactgcaga ggatcgctga gtggggtact gtgtctcata gacatgcgcc     1800

acctccacgt gagaacaagg gtgaaggtga gggaagcccc tcaggttggg tcttttgctg     1860

tgcctcctat gtatgtctgg tttgctggaa gagtgattaa tacatctttta atttattaaa    1920

aaacaatgta gacctttaaa cttcagtctt attgggaata aaagggaact taattcatac     1980

aggtacttga tacagtttata cattttccac ttacaaaag aagacaattc tgttaaatga     2040

aacgtgtatc gtaaaatgta attttattta cccacgagaa tgttgttatt ttagcaatag     2100

aactcaatgc agatgcattg gttattaccc tgtgtgtaccct gtccctcatt ttgctgtgac    2160

accctgaaaa agctgaccac aaatgcagta ttatcattga catacctctg tcc             2213


<210>  74

<211>  2201

<212>  DNA

<213>  NM_005520.1│ Homo sapiens heterogeneous nuclear ribonucleoprotein H1 (H)
(HNRPH1), mRNA


<400>  74
ttttttttttt cgtcttagcc acgcagaagt cgcgtgtcta gtttgtttcg acgccggacc       60
```

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gcgtaagaga cgatgatgtt gggcacggaa ggtggagagg gattcgtggt gaaggtccgg    120

ggcttgccct ggtcttgctc ggccgatgaa gtgcagaggt ttttttctga ctgcaaaatt    180

caaaatgggg ctcaaggtat tcgtttcatc tacaccagag aaggcagacc aagtggcgag    240

gcttttgttg aacttgaatc agaagatgaa gtcaaattgg ccctgaaaaa agacagagaa    300

actatgggac acagatatgt tgaagtattc aagtcaaaca acgttgaaat ggattgggtg    360

ttgaagcata ctggtccaaa tagtcctgac acggccaatg atggctttgt acggcttaga    420

ggacttccct ttggatgtag caaggaagaa attgttcagt tcttctcagg gttggaaatc    480

gtgccaaatg ggataacatt gccggtggac ttccaggga ggagtacggg ggaggccttc    540

gtgcagtttg cttcacagga aatagctgaa aaggctctaa agaaacacaa ggaaagaata    600

gggcacaggt atattgaaat ctttaagagc agtagagctg aagttagaac tcattatgat    660

ccaccacgaa agcttatggc catgcagcgg ccaggtcctt atgcagacc tggggctggt    720

agagggtata acagcattgg cagaggagct ggctttgaga ggatgaggcg tggtgcttat    780

ggtggaggct atggaggcta tgatgattac aatggctata atgatggcta tggatttggg    840

tcagatagat ttggaagaga cctcaattac tgttttttcag gaatgtctga tcacagatac    900

ggggatggtg gctctacttt ccagagcaca acaggacact gtgtacacat gcgggggatta    960

ccttacagag ctactgagaa tgacatttat aatttttttt caccgctcaa ccctgtgaga    1020

gtacacattg aaattggtcc tgatggcaga gtaactggtg aagcagatgt cgagttcgca    1080

actcatgaag atgctgtggc agctatgtca aaagacaaag caaatatgca acacagatat    1140

gtagaactct tcttgaattc tacagcagga gcaagcggtg gtgcttacga cacagatat    1200

gtagaactct tcttgaattc tacagcagga gcaagcggtg gtgcttatgg tagccaaatg    1260

atgggaggca tgggcttgtc aaaaccagtcc agctacgggg gcccagccag ccagcagctg    1320

agtgggggt acggaggcgg ctacggtggc cagagcagca tgagtggata cgaccaagtt    1380

ttacaggaaa actccagtga ttttcaatca aacattgcat aggtaaccaa ggagcagtga    1440

acagcagcta ctacagtagt ggaagccgtg catctatggg cgtgaacgga atgggagggt    1500

tgtctagcat gtccagtatg agtggtggat ggggaatgta attgatcgat cctgatcact    1560

gactcttggt caacctttt ttttttttt ttttctttaa gaaaacttca gtttaacagt    1620

ttctgcaata caagcttgtg atttatgctt actctaagtg gaaatcagga ttgttatgaa    1680

gacttaaggc ccagtatttt tgaatacaat actcatctag gatgtgaacag tgaagctgag    1740
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taaactataa ctgttaaact taagttccag cttttctcaa gttagttata ggatgtactt    1800

aagcagtaag cgtatttagg taaaagcagt tgaattatgt taaatgttgc cctttgccac    1860

gttaaattga acactgtttt ggatgcatgt tgaaagacat gctttttattt ttttttgtaaa    1920

acaatatagg agctgtgtct actattaaaa gtgaaacatt ttggcatgtt tgttaattct    1980

agtttcattt aataacctgt aaggcacgta agtttaagct ttttttttttt ttaagttaat    2040

gggaaaaatt tgagacgcaa taccaatact taggattttg gtcttggtgt ttgtatgaaa    2100

ttctgaggcc ttgatttaaa tctttcattg tattgtgatt tccttttagg tatattgcgc    2160

taagtgaaac ttgtcaaata aatcctcctt ttaaaaactg c                        2201


<210>  75

<211>  1895

<212>  DNA

<213>  NM_004046.4| Homo sapiens ATP synthase, H+ transporting, mitochondrial F1
complex, alpha subunit, isoform 1, cardiac muscle (ATP5A1), nuclear gene
encoding mitochondrial protein, transcript variant 2, mRNA


<400>  75
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agaagtaccg cctgcggagt aactgcaaag atgctgtccg tgcgcgttgc tgcggccgtg     120

gtccgcgccc ttcctcggcg ggccggactg gtctccagaa atgctttggg ttcatctttc     180

attgctgcaa ggaacttcca tgcctctaac actcatcttc aaaagactgg gactgctgag     240

atgtcctcta ttcttgaaga gcgtattctt ggagctgata cctctgttga tcttgaagaa     300

actgggcgtg tcttaagtat tggtgatggt attgcccgcg tacatgggct gaggaatgtt     360

caagcagaag aaatggtaga gttttcttca ggcttaaagg gtatgtcctt gaacttggaa     420

cctgacaatg ttggtgttgt cgtgtttgga aatgataaac taattaagga aggagatata     480

gtgaagagga caggagccat tgtggacgtt ccagttggtg aggagctgtt gggtcgtgta     540

gttgatgccc ttggtaatgc tattgatgga aagggtccaa ttggttccaa gacgcgtagg     600

cgagttggtc tgaaagcccc cggtatcatt cctcgaattt cagtgcggga accaatgcag     660

actggcatta aggctgtgga tagcttggtg ccaattggtc gtggtcagcg tgaactgatt     720

attggtgacc gacagactgg gaaaacctca attgctattg acacaatcat taaccagaaa     780

cgtttcaatg atggatctga tgaaaagaag aagctgtact gtatttatgt tgctattggt     840
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caaaagagat ccactgttgc ccagttggtg aagagactta cagatgcaga tgccatgaag     900

tacaccattg tggtgtcggc tacggcctcg gatgctgccc cacttcagta cctggctcct     960

tactctggct gttccatggg agagtatttt agagacaatg gcaaacatgc tttgatcatc    1020

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agagcagcca aaatgaacga tgcttttggt ggtggctcct tgactgcttt gccagtcata    1200

gaaacacagg ctggtgatgt gtctgcttac attccaacaa atgtcatttc catcactgac    1260

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ctgaagcaag gacagtattc tcccatggct attgaagaac aagtggctgt tatctatgcg    1560

ggtgtaaggg gatatcttga taaactggag cccagcaaga ttacaaagtt tgagaatgct    1620

ttcttgtctc atgtcgtcag ccagcaccaa gccttgttgg gcactatcag ggctgatgga    1680

aagatctcag aacaatcaga tgcaaagctg aaagagattg taacaaattt cttggctgga    1740

tttgaagctt aaactcctgt ggattcacat caaataccag ttcagttttg tcattgttct    1800

agtaaattag ttccatttgt aaaagggtta ctctcatact ccttatgtac agaaatcaca    1860

tgaaaaataa aggttccata atgcatagtt aaaaa                               1895


<210>  76

<211>  1290

<212>  DNA

<213>  NM_001970.3| Homo sapiens eukaryotic translation initiation factor 5A
(EIF5A), mRNA


<400>  76
gcggcggcgg cggtagaggc ggcggcggcg gcggcagcgg gctcggaggc agcggttggg      60

ctcgcggcga gcggacgggg tcgagtcagt gcgttcgcgc gagttggaat cgaagcctct     120

taaaatggca gatgacttgg acttcgagac aggagatgca ggggcctcag ccaccttccc     180

aatgcagtgc tcagcattac gtaagaatgg ctttgtggtg ctcaaaggcc ggccatgtaa     240
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gatcgtcgag atgtctactt cgaagactgg caagcacggc cacgccaagg tccatctggt      300

tggtattgac atctttactg ggaagaaata tgaagatatc tgcccgtcaa ctcataatat      360

ggatgtcccc aacatcaaaa ggaatgactt ccagctgatt ggcatccagg atgggtacct      420

atcactgctc caggacagcg gggaggtacg agaggacctt cgtctccctg agggagacct      480

tggcaaggag attgagcaga agtacgactg tggagaagag atcctgatca cggtgctgtc      540

tgccatgaca gaggaggcag ctgttgcaat caaggccatg gcaaataac tggctcccag       600

gatggcggtg gtggcagcag tgatcctctg aacctgcaga ggccccctcc ccgagcctgg      660

cctggctctg gcccggtcct aagctggact cctcctacac aatttatttg acgttttatt      720

ttggttttcc ccacccctc aatctgtcgg ggagcccctg ccttcacct agctcccttg        780

gccaggagcg agcgaagctg tggccttggt gaagctgccc tcctcttctc ccctcacact      840

acagccctgg tggggggagaa ggggtgggt gctgcttgtg gtttagtctt ttttttttt      900

tttttttttt tttaaattc aatctggaat cagaaagcgg tggattctgg caaatggtcc       960

ttgtgccctc cccactcatc cctggtctgg tccctgttg cccatagccc tttaccctga       1020

gcaccacccc aacagactgg ggaccagccc cctcgcctgc ctgtgtctct cccaaaccc       1080

ctttagatgg ggagggaaga ggaggagagg ggagggggacc tgccccctcc tcaggcatct     1140

gggagggccc tgcccccatg ggctttaccc ttccctgcgg gctctctccc cgacacattt      1200

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aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa                                      1290
```

<210> 77

<211> 2512

<212> DNA

<213> NM_005041.3| Homo sapiens perforin 1 (pore forming protein) (PRF1), mRNA


```
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tcggtgaaga gaggatatcc atctgtgtag ccgcttctct atacgggatt ccagctccat      120

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cccgtgccac acagccgcac gctcagagtg caagcgcagc cacaagttcg tgcctggtgc      240
```

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atggctggcc ggggagggtg tggacgtgac cagcctccgc cgctcgggct ccttcccagt    300

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ccatgtggta cacactcccc cgctgcaccc tgacttcaag agggccctcg gggacctgcc    720

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ggcccaggtc aacataggca tccacggcag catctctgcc gaagccaagg cctgtgagga    960

gaagaagaag aagcacaaga tgacggcctc cttccaccaa acctaccggg agcgccactc    1020

ggaagtggtt ggcggccatc acacctccat taacgacctg ctgttcggga tccaggccgg    1080

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caagtctggt tcccatgagg tgagatgcaa cctgaatcat ggcacctaa aattccgcta    1680

tcatgccagg tgcttgcccc acctgggagg aggcacctgc ctggactatg tcccccaaat    1740

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tattcccatt agaccaagct tgtccaaccc gaggcccgca tgcggcccag gatgcctttg    1920

aatgcggccc aacgcaaatt cgcaaacttt cttaaaacat tatgagtttc tttttgctat    1980
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ctttgccctt gagacaaatg gaggcctgtc ctgacctgtc tcaccatgta catagcttga    2460

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<210>    78

<211>    4623

<212>    DNA

<213>    NM_014965.2| Homo sapiens OGT(O-Glc-NAc transferase)-interacting protein
106 KDa (OIP106), mRNA


<400>    78
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taggggtcg gggcacccc caaggatggt ccttagggt gatgttttgg ctttggggtg     180

acttcagcaa tgtccctgcg agacaagggc ggggaagaag aatgttttga atacgactgc     240

caggatgaag agaggaagcc aacccacagg cagcatgaca cccaggacct cttggaagag     300

gtttttatgtg ctgaaagagt tggccagatg actaagacat ataatgacat agatgctgtc     360

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cacatcaggg aggaggtgtc tcagctccgg catgagctgt ccatgaagga tgagctgctt     540

cagttctaca ccagcgctgc ggaggagagt gagcccgagt ccgtttgctc aaccccgttg     600

aagaggaatg agtcgtcctc ctcagtccag aattactttc atttggattc tcttcaaaag     660

aagctgaaag accttgaaga ggagaatgtt gtacttcgat ccgaggccag ccagctgaag     720

acagagacca tcacctatga ggagaaggag cagcagctgg tcaatgactg cgtgaaggag     780
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aaaaaggcaa aagcttgcgc agtggaaaat gaagaacttg tccagcatct gggggctgct      960

aaggatgccc agcggcagct cacagccgag ctgcgtgagc tggaggacaa gtacgcagag     1020

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cccaatacca cgtctcggcg ctaccactca ctgggcctgt ttcccatgga ttccttggca     1140

gcagagattg agggaacgat gcgcaaggag ctgcagttgg aagaggccga gtctccagac     1200

atcactcacc agaagcgtgt ctttgagaca gtaagaaaca tcaaccaggt tgtcaagcag     1260

agatctctga cccttctcc catgaacatc cccggctcca accagtctc ggccatgaac      1320

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aaaaaaaaaa                                                          4930


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<211>  702

<212>  DNA

<213>  NM_014183.2| Homo sapiens dynein, cytoplasmic, light polypeptide 2A
(DNCL2A), transcript variant 1, mRNA


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atggcagagg tggaggagac actgaagcga ctgcagagcc agaagggagt gcagggaatc     120

atcgtcgtga acacagaaag cattcccatc aagagcacca tggacaaccc caccaccacc     180

cagtatgcca gcctcatgca cagcttcatc ctgaaggcac ggagcaccgt gcgtgacatc     240

gaccccagga acgatcctcac cttccttcga attcgctcta agaaaaatga aattatggtt     300

gcaccagata aagactattt cctgattgtg attcagaatc caaccgaata agccactctc     360

ttggctccct gtgtcattcc ttaatttaat gcccccaag aatgttaatg tcaatcatgt     420

cagtggacta gcacatggca gtcgcttgga acccactcac accaatccag tgaccgtgtg     480
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acccagcttc cttctgacct tcagttcact ttgtcgccct tggagaaagc tgttttttctt     660

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<211> 2100

<212> DNA

<213> NM_015907.2| Homo sapiens leucine aminopeptidase 3 (LAP3), mRNA


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ccggctgcgg ggcgagtagt cgtccgacgt ctggccgtga gacgtttcgg gagccggagt      240

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caaaaaaaga agatggctgt gtcggcaaag ctctatggaa gtgggatca ggaggcctgg      720

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gccaatgaga tgacgccaac cagatttgct gaaattattg agaagaatct caaaagtgct     840

agtagtaaaa ccgaggtcca tatcagaccc aagtcttgga ttgaggaaca ggcaatggga     900

tcattcctca gtgtggccaa aggatctgac gagccccag tcttcttgga aattcactac      960

aaaggcagcc ccaatgcaaa cgaaccaccc ctggtgtttg ttgggaaagg aattaccttt     1020

gacagtggtg gtatctccat caaggcttct gcaaatatgg acctcatgag ggctgacatg     1080
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gccttgattt tttttcatt tcacacaaag atttataaag gtaaagttaa tatcttactt     1920

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<211>  1510

<212>  DNA

<213>  NM_018478.1| Homo sapiens chromosome 20 open reading frame 35 (C20orf35),
mRNA


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gtctgcgcgg ggcggggcg cagcaagtgc atccgagcga gcggagacta gcgcaccggc     180

gtcggtggcg agggtggtgc agaggagtcc ggctgggcgg agggaggaag gatgggtgcg    240

ggtaactttt tgaccgcctt ggaagtacca gtagccgcgc tcgcaggggc tgcctccgac    300
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aaaaaaaaaa                                                         1510


<210>  86

<211>  3105

<212>  DNA

<213>  NM_030674.2| Homo sapiens solute carrier family 38, member 1 (SLC38A1),
mRNA


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ggactaattt tcaaactaca gacactttct aatcatgatg catttcaaaa gtggactcga   480

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ctaggttatt acaaaatacaa gaattttgta ccttaaataa gcctcatttc tatttcttct   3060

tcattaattc tccatctagt cttgtgaaaa aaaaaaaaaa aaaaa           3105
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<210> 87

<211> 2711

<212> DNA

<213> NM_016028.4| Homo sapiens suppressor of variegation 4-20 homolog 1
(Drosophila) (SUV420H1), transcript variant 2, mRNA

<400> 87
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tgtcagctct aacactgatg cagataccac tcaggaaaaa aacaatgcaa gtaagtaagg   1440

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cagcagacca ccagtaccag actcagaatt gtgatagagg agcattttga acagtgccgt   1980

attgtgacat gctgtattgg ctactccaga aagtaggagt aaagatggaa aggagaaaga   2040

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<211>  2977

<212>  DNA

<213>  NM_022105.2| Homo sapiens death associated transcription factor 1
(DATF1), transcript variant 1, mRNA


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cctcccggaa gcgcggaacc tcagcttccg tacttgcgca gaactcccct cgcggcgacc    180

acgcactacg ggttggcgcc agagtcaaaa ggcgtcggcc ctctggcaag atggctgctg    240
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gcctgttact cgtgaacagt ggctgacaac agtgttgttg tgagcctggc tgtctgcttg      420

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cccacagcag cagctgggcc tgtccctgcg gcgcagtggg aggcagccca agcgcactga      720

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agaaaaaaaa gagaccacag tgaagaaggc agtggtggtc cctgcgcgga gtgaagcact     1980
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atttcaaaca gatctttgaa acctcagatt ctgtggtgca attttaatgt tttcttgttt       2940

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<211> 1047

<212> DNA

<213> NM_018487.2| Homo sapiens hepatocellular carcinoma-associated antigen 112 (HCA112), mRNA

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gtggcctcgt gggtgatgca gatcgtgctg gggatcttga gtgcagtcct aggaggattt      300
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catgttattg ttcccaggaa aaaaaaa                                        1047


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<213>  NM_014454.1| Homo sapiens sestrin 1 (SESN1), mRNA


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agtatcttcg ttcggtgaaa gaaacacctc atcgtccatc agacgggctt tcaaataccg       180

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ccttcaaaga tgtgagagag aaaagtgagt ttattctgaa gagcatccag gaacttggca       300

ttagaattcc tcgaccacta ggacagggac caagcagatt catcccagaa aaggagatcc       360

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ctcaaaaact acagaattta ggagaactta acaaagtgtt agcccataga ccttggctta      720

ttaccaaaga acacattgag ggactttaa aagctgaaga gcacagctgg tcccttgcgg       780

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aataagataa caagaattgg agcatgcaaa gaatgggact tggataatga cttaagcttt     2040

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acataacttg tgtgatttca aattaaactt gcattatgtg ttaattttct tgcatctaaa     2220

aaagcataga attcctactc acacagctca gcaacaacca ttttgatggt aacagttaat     2280
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<211>  3802

<212>  DNA

<213>  NM_017763.1| Homo sapiens hypothetical protein FLJ20315 (FLJ20315), mRNA


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<211>  1236

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tgaaggaaca gctgcagccc cttgaacagg tgaaagctgg aatagaagct cattcggaag     720

ccaaaaccag tggactcctg tgggctggat tggcactgct gtccattcag ggtggggcac     780

tggcctggct cacgtggtgg gtgtactcct gggatatcat ggagccagtt acatacttca     840

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agaattaatc ttacagtttt aaatgtcgtc agattttcca ttatgtattg attttgcaac    1140

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<210>  93

<211>  2096

<212>  DNA

<213>  NM_024792.1| Homo sapiens membrane protein expressed in epithelial-like
lung adenocarcinoma (CT120), mRNA


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aagactgtgc taccatgtgt tctcaagtcg tagtttaaaa agtggatttt taaagtgcct    2040

ttcaattgtc tgtgaacgtc taaaggactg atttgtctca aaaaaaaaaa aaaaaa        2096


<210>  94

<211>  4372

<212>  DNA

<213>  NM_014314.2| Homo sapiens DEAD (Asp-Glu-Ala-Asp) box polypeptide 58
(DDX58), mRNA


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cctgcgggga acgtagctag ctgcaagcag aggccggcat gaccaccgag cagcgacgca     180

gcctgcaagc cttccaggat tatatccgga agaccctgga ccctacctac atcctgagct     240

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ccattgaaag ttgggatttc aaaaaaaattg aaaagttgga ggagtataga ttactttttaa   480

aacgtttaca accagaattt aaaaaccagaa ttatcccaac cgatatcatt tctgatctgt    540

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ccaaaacttt gaaacttgct ttggagaaag aaaggaacaa gttcagtgaa ctgtggattg    720

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acatacagat tttctaccaa gaagatccag aatgccagaa tcttagtgag aattcatgtc    840

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tgctctgcag aaagtgcaaa gccttggcat gttacacagc tgacgtaaga gtgatagagg    2640

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cataaagcca tttttatgct tactactaaa agctgtccac tgtagagttg ctgtatgtag    3720

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taagacccaa tacaaatgga ttatttaaac caaaaaaaaa aaaaaaaaaa aa           4372
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<210> 95

<211> 2163

<212> DNA

<213> NM_015515.3| Homo sapiens keratin 23 (histone deacetylase inducible) (KRT23), transcript variant 1, mRNA

<400> 95
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ctaatcaaag tgcgagttta tgagggtaaa gctctacttt cctactgcag ccttcagatt    2100

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aaa                                                                  2163


<210> 96

<211> 2881

<212> DNA

<213> NM_007210.2| Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide
N-acetylgalactosaminyltransferase 6 (GalNAc-T6) (GALNT6), mRNA


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a                                                                    2881


<210>  97

<211>  1930

<212>  DNA

<213>  NM_020183.3| Homo sapiens aryl hydrocarbon receptor nuclear translocator-
like 2 (ARNTL2), mRNA


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agtgattcag acccatccca gtcaggaatc atgacagaaa aagtggtgga aaagctttct     240

cagaatcccc ttacctatct tctttcaaca aggatagaaa tatcagcctc cagtggcagc     300

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<211> 3394

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<211>  2524

<212>  DNA

<213>  NM_000249.2| Homo sapiens mutL homolog 1, colon cancer, nonpolyposis type
2 (E. coli) (MLH1), mRNA


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<211> 3262

<212> DNA

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<400> 110

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ataccatgtg ccgctgtgtc accaaacaga tccaaatgtg agagaaacac agtggcagat      780

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<211> 3138

<212> DNA

<213> NM_018223.1| Homo sapiens checkpoint with forkhead and ring finger domains (CHFR), mRNA

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<211> 2466

<212> DNA

<213> NM_024645.1| Homo sapiens hypothetical protein FLJ13842 (FLJ13842), mRNA

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<211> 3658

<212> DNA

<213> NM_025195.2| Homo sapiens tribbles homolog 1 (Drosophila) (TRIB1), mRNA

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mRNA


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tgtcttaccc ccctgggaca gctgttacct ttgcagtgtt gccgaatcac agcagttacc   1080

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<211>  2152

<212>  DNA

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transcript variant 2, mRNA


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tggattttct ctacaaaaat tattgagcaa ccctaattaa cctgattttt tgctgataat    180

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tttttaaatt taagaataag acacacaaaa ctctattaaa aaaaaaaaag aaataaacct      2040

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<211>  3010

<212>  DNA

<213>  NM_145343.1| Homo sapiens apolipoprotein L, 1 (APOL1), transcript variant
2, mRNA



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tctgccaggg gaagattcct tgacttctgg ggtgatggaa aagaaacagg ctgtgctgtg        180

tccctaatgg gaaacgtggc tgagacaggg gagtgagaag ggtgcgttgc agaatggtgc        240
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<211>  2759

<212>  DNA

<213>  NM_080796.1| Homo sapiens death associated transcription factor 1
(DATF1), transcript variant 2, mRNA


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(DNCL2A), transcript variant 2, mRNA

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tgcaccaacc ttcccagag ctccggagcg ccctctcctc acttccaggt tttggagcaa      660

gagcttgcag gaagcccgca cccagcttcc ttctgacctt cagttcactt tgtcgccctt    720

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(STAU), transcript variant T3, mRNA


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member 6 (TNFRSF6), transcript variant 2, mRNA


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<210> 130

<211> 2730

<212> DNA

<213> NM_152872.1| Homo sapiens tumor necrosis factor receptor superfamily, member 6 (TNFRSF6), transcript variant 3, mRNA

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ttacgagtga cttggctgga gcctcagggg cgggcactgg cacggaacac accctgaggc     240

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attgtgtgat gaaggacatg gcttagaagt ggaaataaac tgcacccgga cccagaatac     720

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<210>  131

<211>  2563

<212>  DNA

<213>  NM_152874.1| Homo sapiens tumor necrosis factor receptor superfamily,
member 6 (TNFRSF6), transcript variant 8, mRNA
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<400> 131

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tcgaaagaat ggtgtcaatg aagccaaaat agatgagatc aagaatgaca atgtccaaga    960

cacagcagaa cagaaagttc aactgcttcg taattggcat caacttcatg gaaagaaaga   1020

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acctaaagaa cttccattta tggaggattt ttttgcccct tgtgtttgga attataaaat    2460

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aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaa                      2563


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<211>  2445

<212>  DNA

<213>  NM_152876.1| Homo sapiens tumor necrosis factor receptor superfamily,
member 6 (TNFRSF6), transcript variant 6, mRNA


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gaagcgtatg acacattgat taaagatctc aaaaaagcca atctttgtac tcttgcagag    960

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<210>  133

<211>  2508

<212>  DNA

<213>  NM_152877.1| Homo sapiens tumor necrosis factor receptor superfamily,
member 6 (TNFRSF6), transcript variant 7, mRNA


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aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaa              2508


<210>   134

<211>   2583

<212>   DNA
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<213> NM_152875.1| Homo sapiens tumor necrosis factor receptor superfamily,
member 6 (TNFRSF6), transcript variant 5, mRNA

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protein mRNA, complete cds


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atgcacctcc tgcaagaaga gctgctgctc ctgttgcccc ctgggctgtg ccaagtgtgc     180
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ccagggctgc atccgcaaag gggcttcgga aaagtgcagc tgctgtgcct gatgtcggga   240

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<211> 3145

<212> DNA

<213> NM_000251. Homo sapiens mutS...[gi:4557760]

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atatttagta atatttttact ttgaggacat tttcaaagat ttttattttg aaaaatgaga    3060

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aatattgatc taaagcttaa ggactatgga gtggatctta ttgaagtttc agacaatgga     240

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gaagagtacg gtttggagctg ttcggatgct ctgcataatc tttttttacat ctcaggtttc     840

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<213>  NM_000179. Homo sapiens mutS...[gi:4504190]


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